43

## SEQUENCE LISTING

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-:110 · KU, Ming-Qun
         EVANS, Thomas C.
   -:120 - Intein Mediated Peptide Ligation
* :130 - NEB-150PUS
  -:140 - 69/786,009
1 (141 - 1999-09-30
   < 150 - 60 / 102,413
   -1151 - 1999-09-30
  -:150 - PCT/US33/22776
  -:151: 1999-09-30
  -K160 × 3
   -1100 - PatentIn Ver. 2.0
   -.210 - 1
   -011 - 43
   +213 - Artificial Sequence
   -127 O. +
    213 Description of Artificial Sequence: the modified
         C-terminal splice junction of the intein from the
         gyrA gene of Mycobacterium xenopi
   - [4 (H)) - 1
   491.1091.cag ccacgetact ggeoteaceg gttgatager gea
  H2108-2
   00110-39
-2120-DNA
   H213 - Artificial Sequence
   +2250 Description of Artificial Sequence: the
         simplementary strand of the C-terminal splice
         junction of the modified intein from the gyrA
         gove of Mycobacterium xonopi
    -----
   gotar rawincinggrigaggidda atalidatagu tgwungwa
   1210 - 3
   1311 - 69
   <212 - DNA
   <213 - Artificial Sequence
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-1220 ·
+12.13 - Description of Artificial Sequence: the polylinker
       sequence inserted upstream of the modified intoin
       from the gyrA gene of Mycobacterium xenopi
<400.8
togantolag acatatgged atgggtggeg geogeologa gggclotted tgcalcacgg 60
dagaigea
-1210.4
-42211 + 69
+2127 DMA
-MIG - Artificial Sequence
-. 11.
+213 - At position 41, "H" = A or C or T.
· 11.10
+2230 Description of Artificial Sequence: the
      complementary strand of the polylinker inserted
      upstream of the modified intein from the gyrA
      gene of Mycobacterium xenopi
+ 4 (i) + 4
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igtotagat
3.11 Oct 5
- 2011: 6:09
HILLIA DNA
+:213: Artificial Sequence
+02.739 \cdot \texttt{Description of Artificial Sequence: pTXB1 plasmid}
      sequence containing the modified intein from the
      gyrA gene of Mycobacterium xenopi
waktaegtea ggtggeaett ttegggggaaa tgtgegegga aeeestatit gtttattttt 60
omuaatadat toaaaatatgt atcogotoat gagadaataa odotgataaa tgottoaata 120
anatitoogt glogoccol a litecettiit 190
tyrgycallt tycottooty titttyctba obcagaaang niggiyaaag baaaagaiyo 240
ng amandag tigggignan gagiggita batogaabig gaidibaana goggiaagai 300
 softgadagt obtogoodd aagaacgottu tussaatgatg aqualittia aagttistyst \mathbb{R}^{n}
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Egoaggabba officigoret oggeesties gyetggetgg titattgetg ataaatstyr A \, {\mathbb N}
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dit mintaga ahadahadat cachqathaa qoathqahaa chqhcadacc aaghthachd 1020.
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                                                                       €569
 tttgctgaaa ggaggaacta tatccggat
 <210 - 6
 \pm 211 \pm 30
 1212 - PRT
 (213 · Artificial Sequence
 1220 -
  (223) Description of Artificial Sequence: synthetic
       peptide
  Ovs Ala Tyr bys Thr Thr Gin Ala Abn bys His Ile Tie Mai Ala Myd
 Gru Gry Ash Pro Tyr Val Fro Val His The Asp Ara Cor Lai
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-1.310F-7 .111: 13 -1.1121- FF.T -213 - Artificial Sequence -.2201-+2223: Description of Artificial Sequence: the amino acid sequence deduced from the polylinker region of pTXB1 -14001-7 Met Ala Met Gly Gly Gly Arg Leu Glu Gly Ser Ser Cys +.210.+ € -12110-42 HIZIZI INA 3213 · Artificial Sequence -12250-#2230 Lescription of Artificial Sequence: polylinker region upstream of the modified intein from the dyrA gene of Mycobacterium xenopi in pTXB1 101 - 8 42 intanggoda tgggtggegg degectegag ggetetteet ge 12100 9  $\pm 1.111 \pm 7$ +:312"+ FFT H213 - Artificial Sequence H220 H H123 - Description of Artificial Sequence: synthetic peptide -1400 - 9 Cys Asp Pro Glu Lys Asp Ser